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## CORRIGENDUM

### Genetic variation in microRNA networks: the implications for cancer research

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There were several mistakes in TABLES 1–3 of this article. In TABLE 1 on page 385, for rs2910164 in hepatocellular carcinoma the odds ratio data refer to males only. For rs11614913 in lung cancer the odds ratio is for both sexes and not for males only. For rs11614913 in oesophageal cancer the reference genotype should be CC/CT and the analysed genotype should be TT. For rs11614913 in breast cancer the reference genotype should be CC and the analysed genotypes should be CT and then TT. For rs895819 in breast cancer the reference genotype should be CT. In TABLE 2 on page 396, for the *SED8* alleles, the miRNA binds the derived alleles and not the ancestral alleles. For the *BCTRP* allele, the miRNA binds deletion alleles that are derived and not insertion alleles that are ancestral. For the *BMPIB* allele, the miRNA binds the C allele and not the derived allele. For the *CD86* allele, the miRNA binds the ancestral allele and not the derived allele. In TABLE 3 on page 398, the odds ratio given for *GEMIN4* rs7813 is for all cases of renal cell carcinoma, not just clear cell carcinoma as indicated. For *GEMIN3* rs197414 the cancer site should be bladder and not renal cell carcinoma and the reference is 35 and not 111. This has been corrected online.